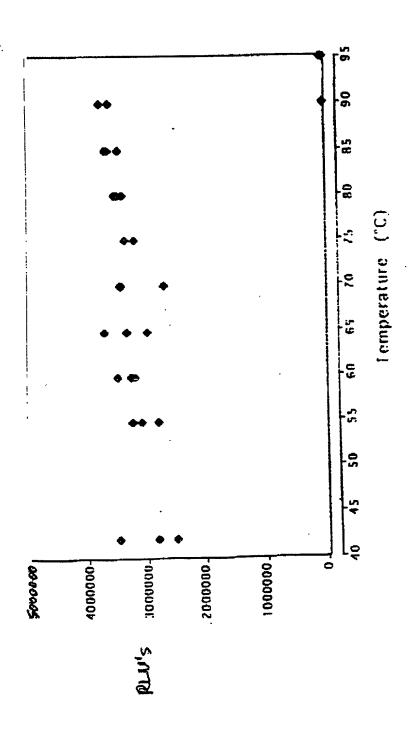
Thermal Stability of Pethets our wide ronge of temperature

CT Amplification after 10 minutes at Each Temperature (10 µL HPA)



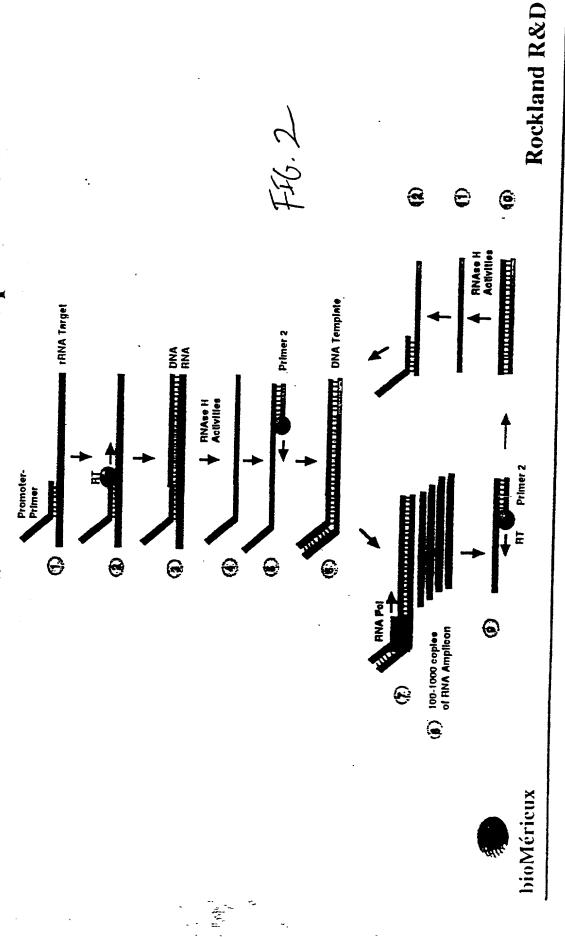
1

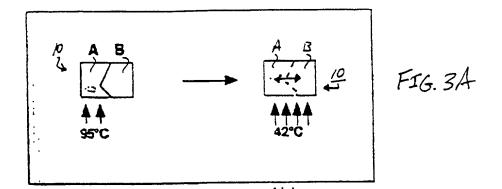
F36.1

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Transcription Mediated Amplification





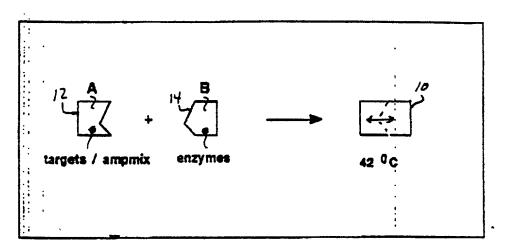
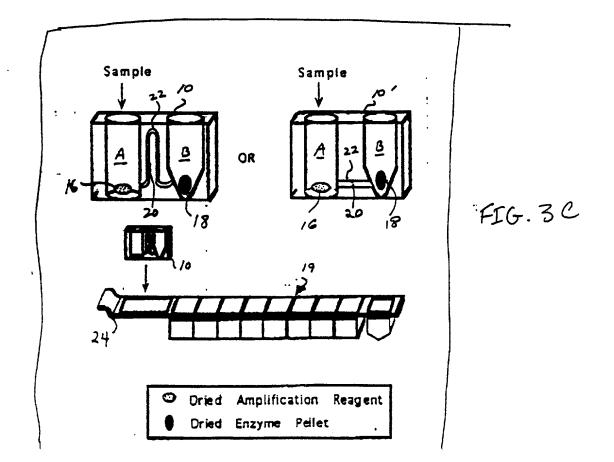
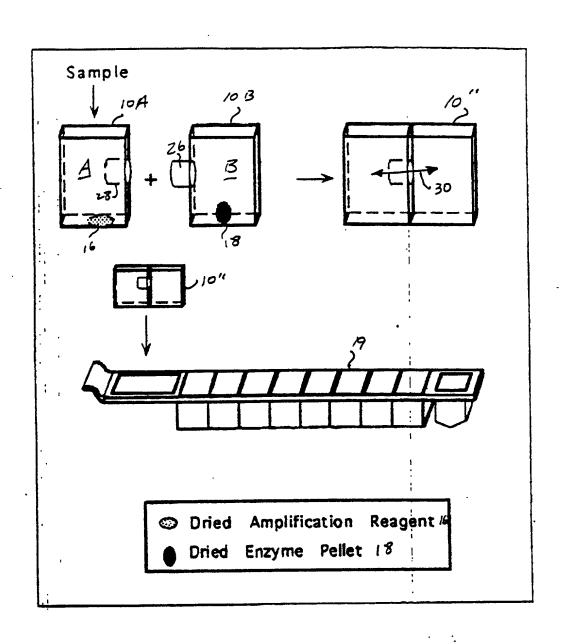
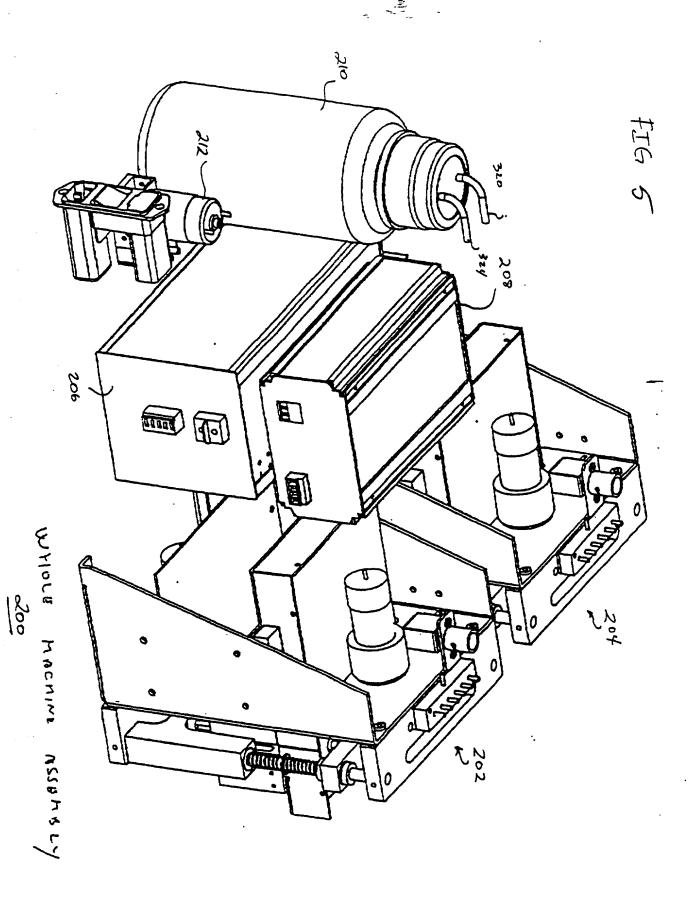


FIG.38





F16.4



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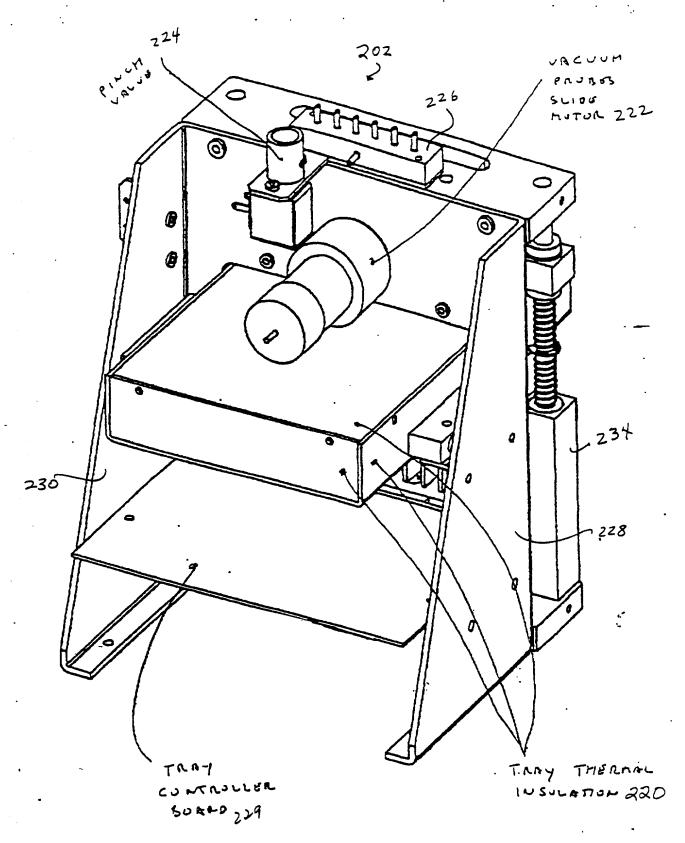


FIG. 6

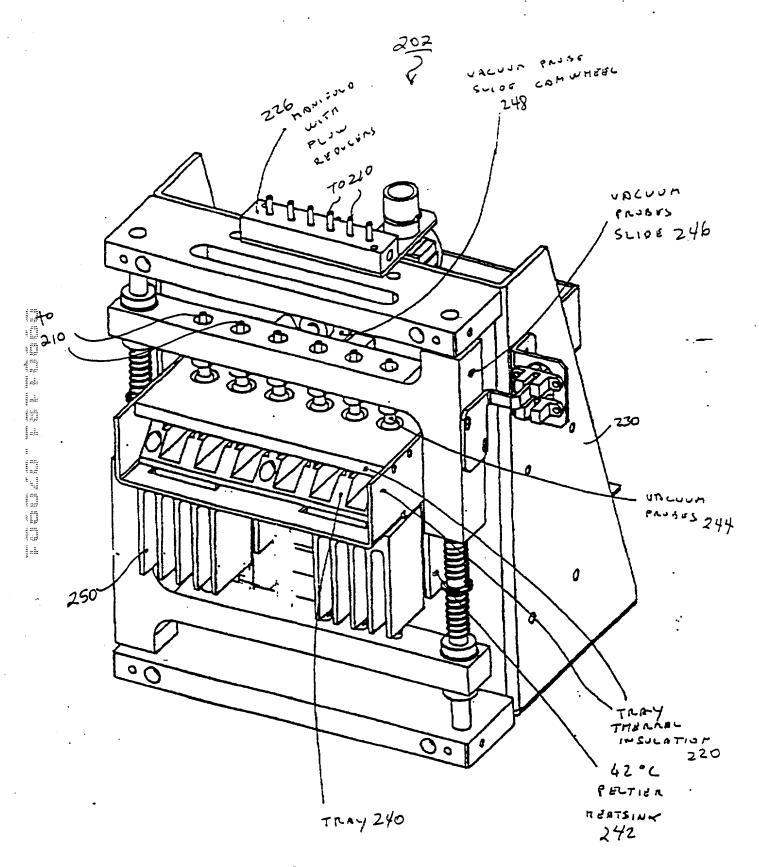
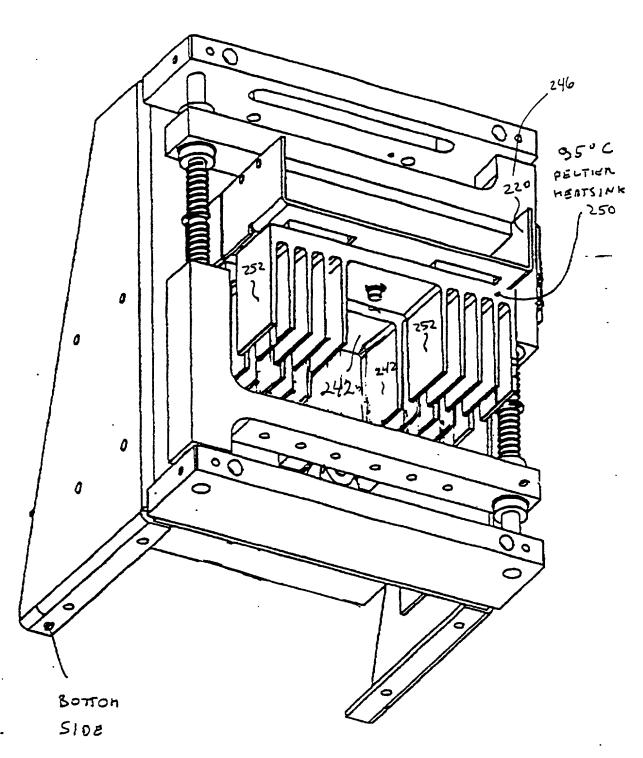


FIG. 7

FIG. 8



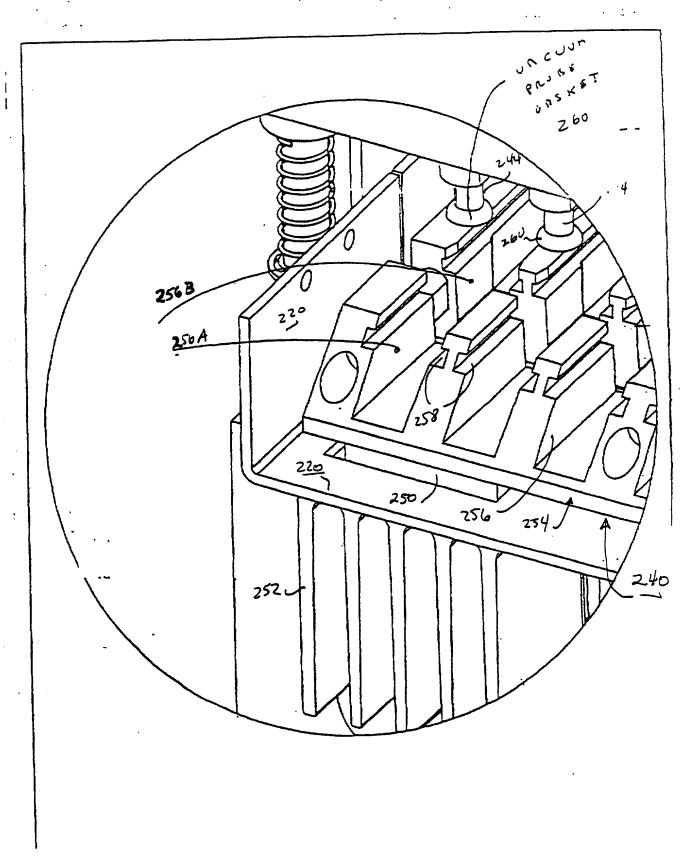
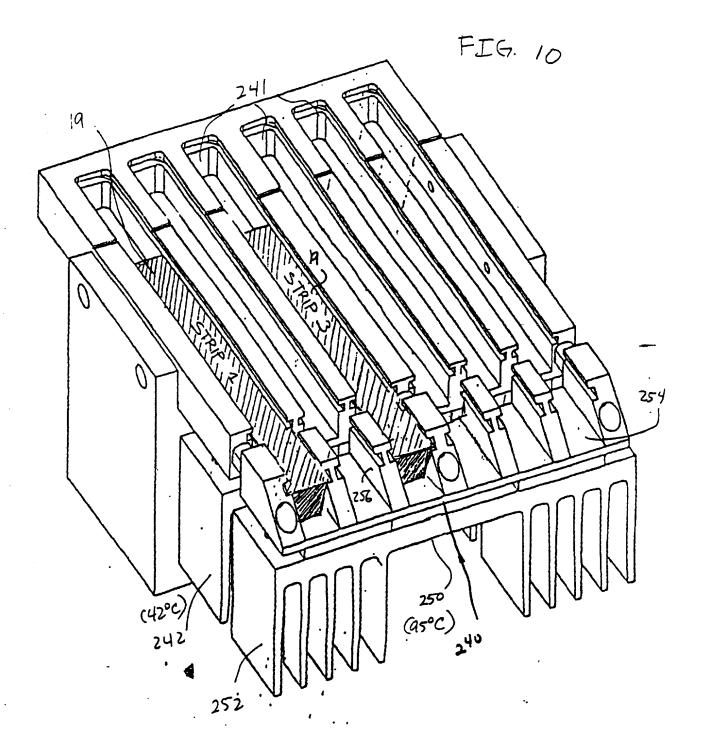


FIG. 9



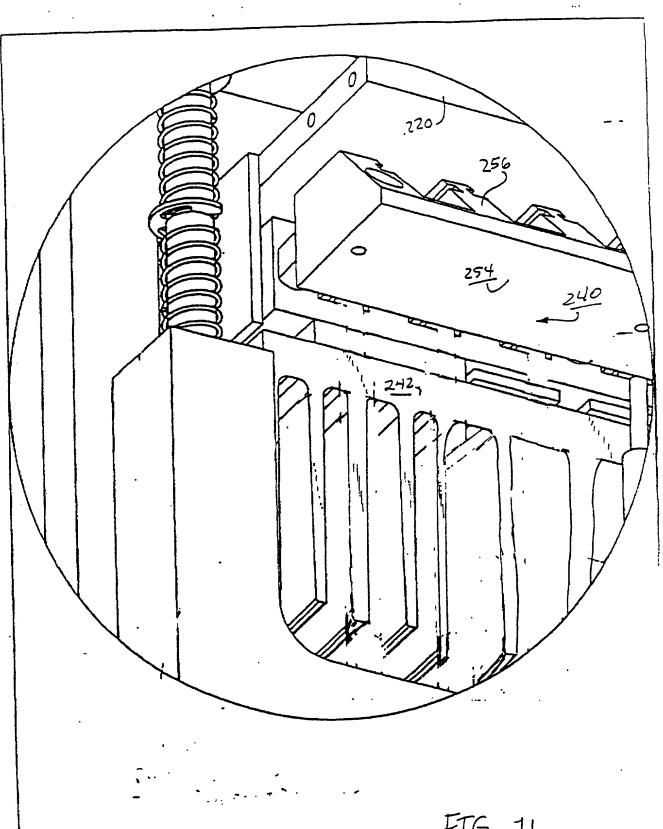
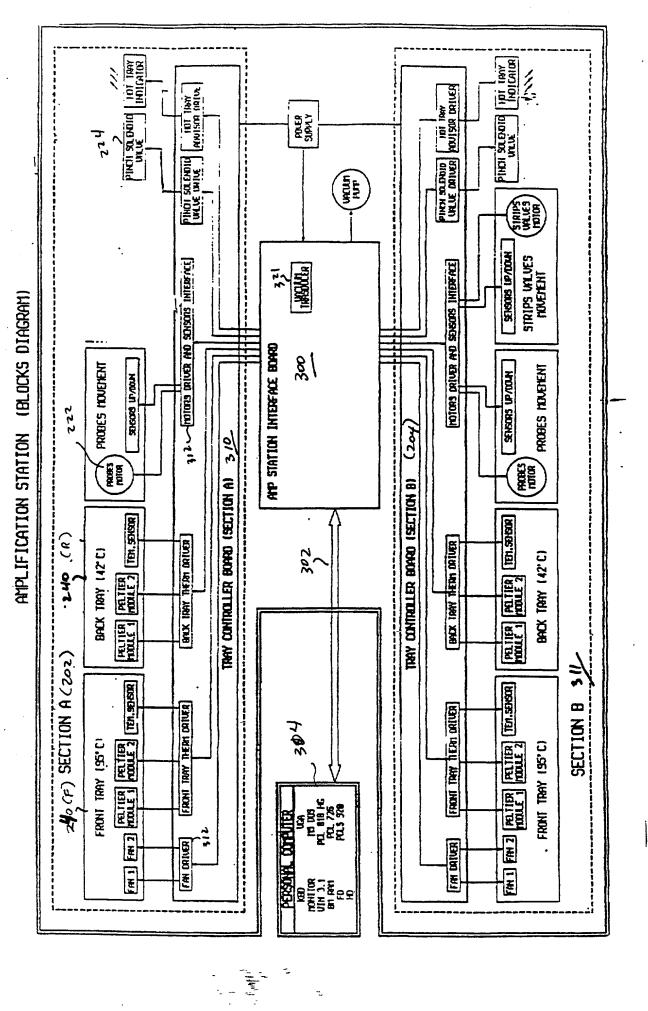
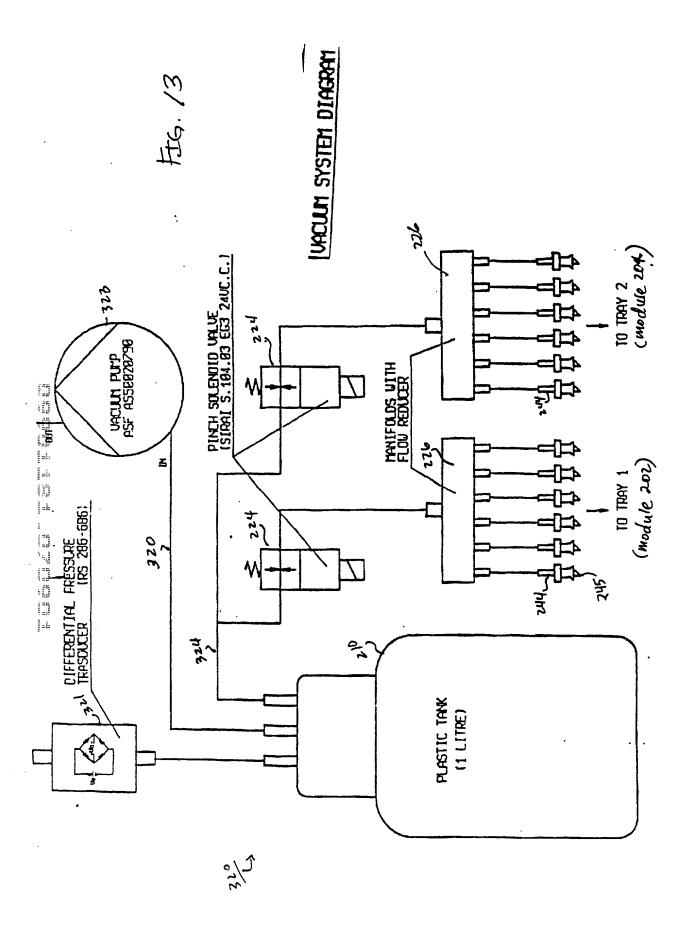


FIG. 11



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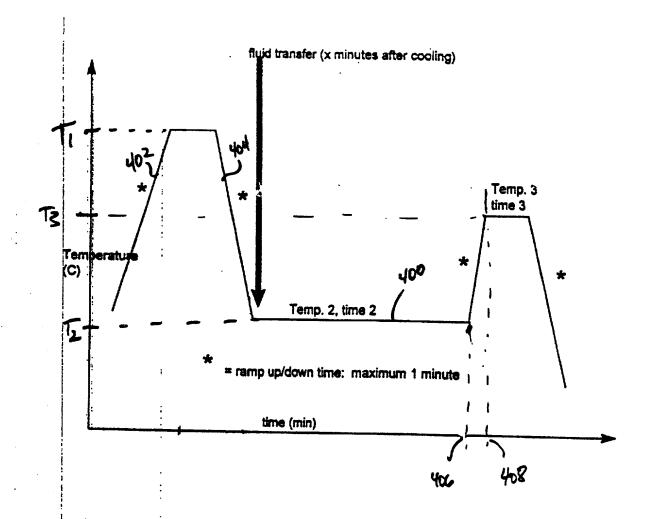


FIG. 14

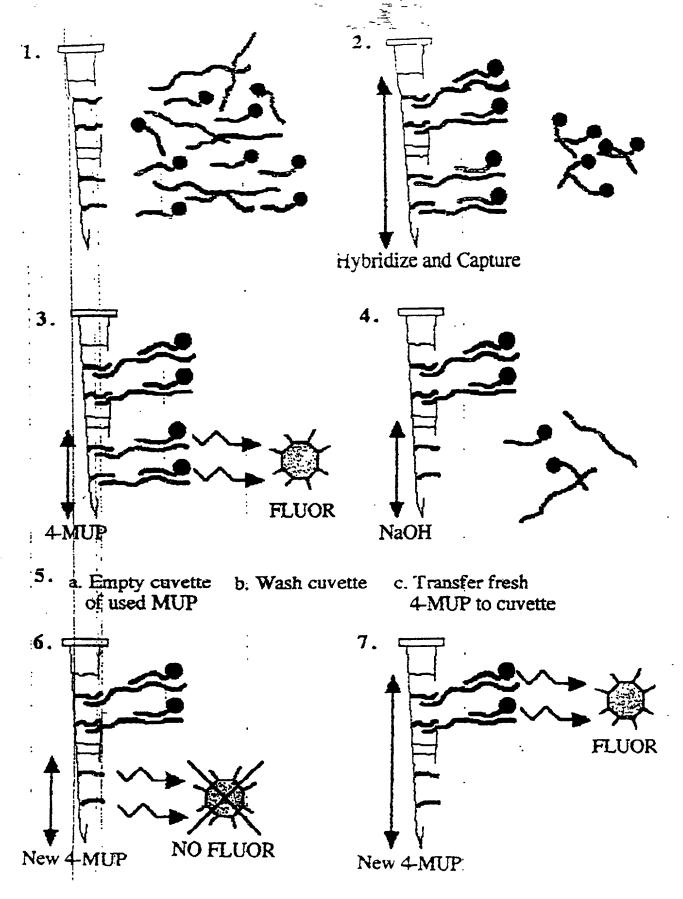
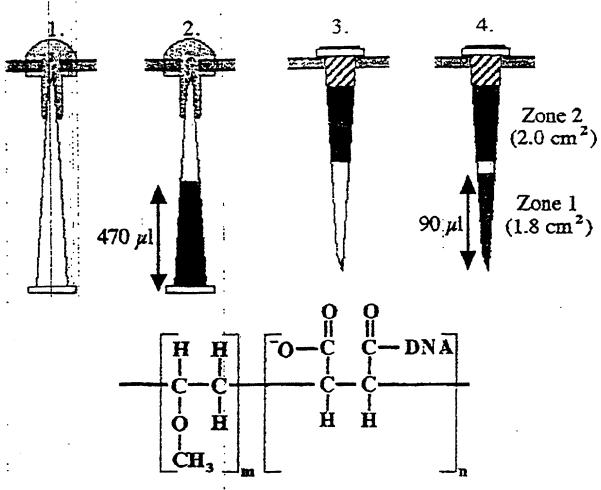


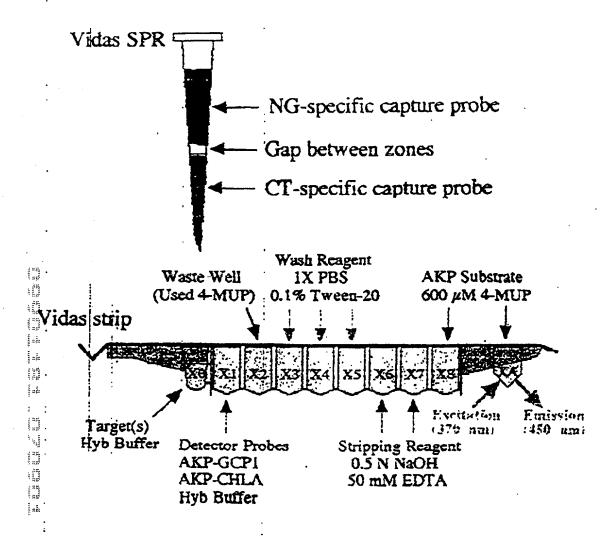
Fig 15

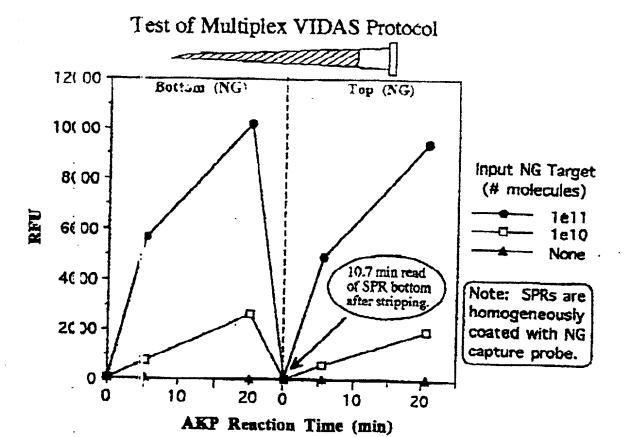
SPR PRODUCTION WITH DISTINCT CAPTURE ZONES



Conjugate of AMVE copolymer and specific capture probe

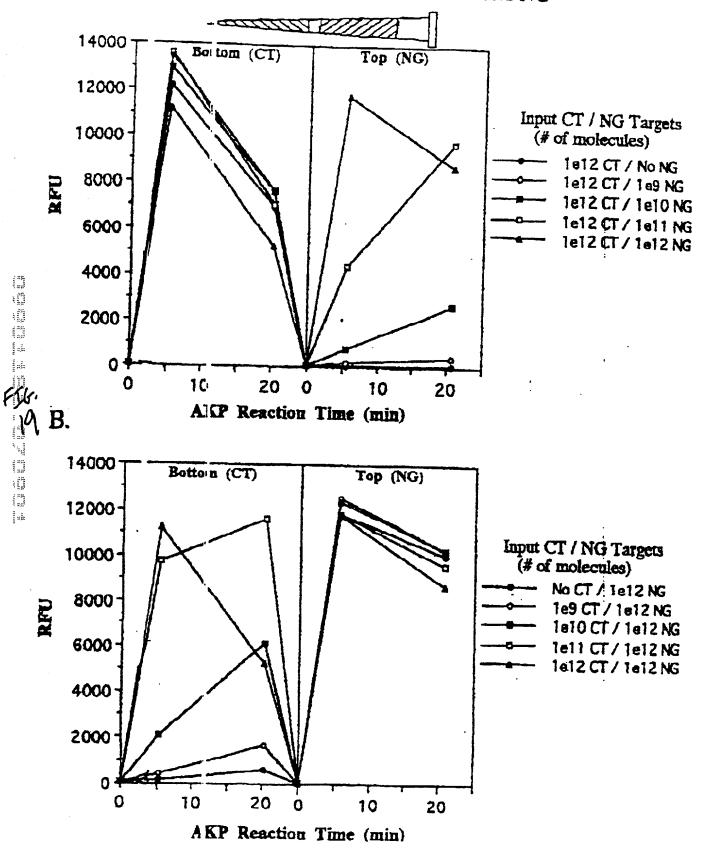
MULTIPLEX STRIP CONFIGURATION



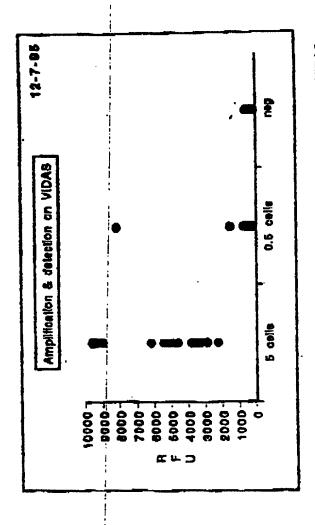


F16-18

FIG. 19 A. Dose Response Multiplex Detection of CT and NG



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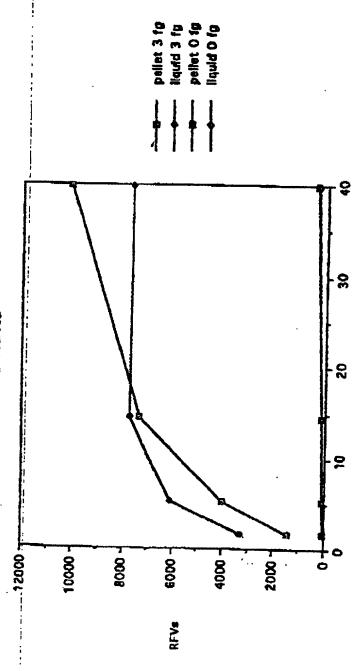


* sample heated off-line in the presence of ampmbs, enzyme transfer by VIDAS

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Binary: VIDAS detection

Jinary: amplification and detection on 44°C VIDAS



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Rend Time

FIG. 22

Rockland R&D

FI6: 23

FIG. 24

Random Internal Control

3'-ccctcgcttacaatcccgtgtgagtacccactcgttcagaaagacattcccgactacagtccgcataactgttcgtactgcttgtt-5'

RAN16 primer: 5'-Agcgaatgttagggcacactc-3' TARGET

RANJ3 AKP-probe: 5'-ATGGGTGAGCAAGTCTTTCTG-3'

3 - - AACTGTTCGTACTGCTGGTCT

(T7 promoter / RAN19 primer) Agaggatatcactcagcataatttaa-5 '

3'-gtacccactcgttcagaaagacattcccgactacagtccgcataactgttcgtactgctggtct-5'

5'-GCAATTAACCCTCACTAAAGGGAGCGAATGTTAGGGCACACTCATGGGTGAGCAAGTC-3'

(T3 promoter)

5'-TCT 99T C9T CAT 9CT T9T CAA TAC 9CC T9A CAT CA9 CCC TTA CA9 AAA 9AC TT9 CTC ACC CAT 9-3'

5'-GCA ATT AAC CCT CAC TAA AGG GAG CGA ATG TTA 999 CAC ACT CAT 999 TGA 9CA AGT C-3'

5'-CAA TAC GCC TGA CAT CAG CCC TTA CAG AAA GAC TTG CTC ACC CAT GAG-3' 5'-AAT TTA ATA CGA CTC ACT ATA 999 AGA TCT 99T CGT CAT 9CT TGT CAA-3'

RIC1 Detection oligo:

RIC1 bottom oligo:

RIC1 top oligo:

T7/RAN19 TMA primer:

RAN21 AMVE-probe: RAN16 TMA primer:

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RAN33 AKP-probet

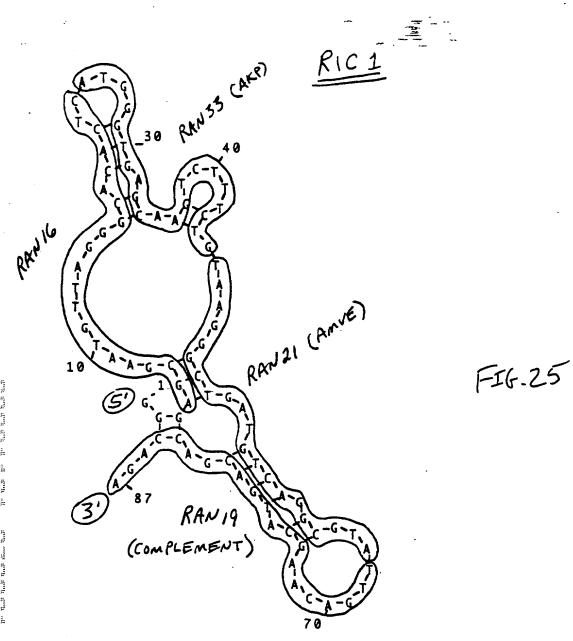
5 -- aminolink-ATg ggT gAg CAA gTC TTT CTg-3' 5 -- aminolink-TAA ggg CTg ATg TCA ggC gTA-3'

5'-AgC GAA TGT TAG GGC ACA CTC-3'

5'-gggägcgaatgttagggcacactcatgggtgägcaaagtctttctgtaagggctgatgtcaggcgtattgacaagcatgacgaccaga-3' 5'-TAAGGGCTGATGTCAGGCGTA-3' RAN21 AMVE-probe

5'-gggagcgaatgttagggcacactcatgggtgagcaaatctttctgtaagggctgatgtcaggcgtattgacaagcatgacgaccaga-3'

The first the first than the train that the train the train that the train that the train that the train than t norm arms, or nates, or owns, or a norm of the second seco



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Random Internal Control 2

5.-cagtagaggtaggggctgctaggagtataacagaagccagtgtacggaacgactcagcacggcgaatactttgctaccagacctagaggagtgcgt-3'

5'-ACGACTCAGCACGGCGAATAC-3' RAN32 AKP-probe RAN51 TMA primer 5'-CAGTAGAGGTAGGGGCTGCTAGGAGT-3'

5 · - CAGTAGAGGTAGGGGGTGCTAGGAGTATAACAGAAGCCAGTGTACGGAACGACTCAGCACGGCGAATACTTTGCTACCAGACCTAGAGGAGTGCGT-3 · TARGET

RAN27 AMVE-probe 5'-TAACAGAAGCCAGTGTACGGA-3'

3'-ACGATGGTCTGGATCTCCTCACGCA

Agaggatatcactcagcataatttaa-5' (T7 promoter / RAN39 primer)

(T3 promoter)

5'-GCAATTAACCCTCACTAAAGGGCAGTAGAGGTAGGGGCTGCTAGGAGTATAACAGAAGCCAGTGTAC-3'

3'-grcttcggtcacatgccttgctgagtcgtgccttatgaaacgatggtctggatctcctcacgca-

OLIGOS TO ORDER!

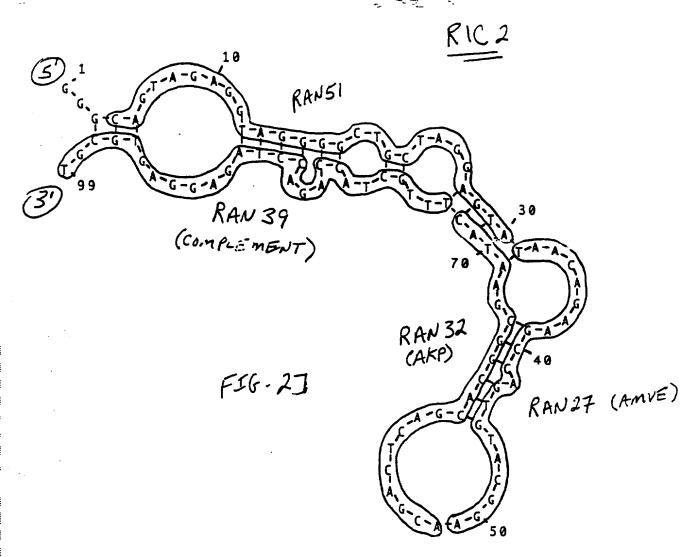
5'-CAG TAG AGG TAG GGG CTG CTA GGA GT-3' RAN51 TWA primer: 5'-aminolink-TAA CAg AAg CCA grg TAC ggA-3' RAN27 AMVB-probes

5'-aminolink-ACG ACT CAG CAC ggc gAA TAC-3' RAN32 AKP-probes

5'-AAT TTA ATA CGA CTC ACT ATA 999 AGA ACG CAC TCC TCT A99 TCT 99T A9C A-3' T7 / RAN39 primer:

5 .- AAG TAT TCG CCG TGC TGA GTC GTT CCG TAC ACT GGC TTC TGT TAT AC-3' RIC2 Detection oligo: 5'-GCA AIT AAC CCT CAC TAA AGG GCA GTA GAG GTA G9G GCT GCT AGG AGT ATA ACA GAA GCC AGT GTA C-3' ic2 Top oligo:

gcc grg crg agr cgr rcc gra cac rgg crr crg-3' 5.-ACG CAC TCC TCT AGG TCT GGT AGC AAA GTA TTC RIC2 Bottom oligo:



Detection of RIC1 DNA Oligo Targets

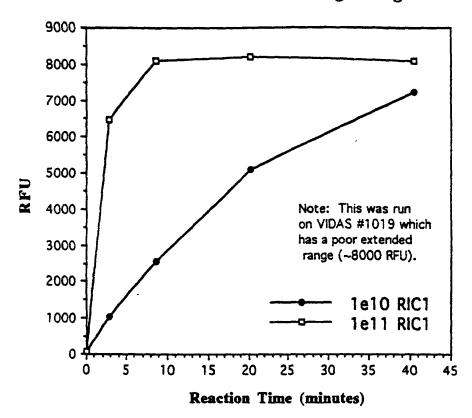


FIG. 28

Detection by 7	Detection by 71 11 protocol		4		-		1	•
B. Kluttz data	OLU PIOLOCOI				• !			
								;
			AKP Type &					
Position	RIC1 RNA*	CT RNA	SPR Type	0 min	1.8 min	5.4 min	14.6 min	40.0 min
5	none	none	RIC1	26	56	58	19	0.2
C2	=	none	RIC1	25	. 55	57	59	99
3	0.1	none	RC1	26	55	22	19	7
2	=	none	RC1	57	26	57	61	89
CS	-	none		56	52	59	65	8
93	=	none	RC1	26	55	57	62	7
10	10	none	RIC1	55	78	114	202	414
D2	=	none	RC1	26	56	59	99	82
D3	100	none	RIC1	26	55	58	62	2
04	=	none	RIC1	22	57	19	70	94
D5	1000	none	RIC1	56	58	81	119	227
90	=	none	RC1	56	22	02	102	184
E	10000	none	RIC1	26	93	500	414	948
E2	2	none	RIC1	56	105	246	497	115
E3	100000	none	RIC1	26	395	1474	3029	6510
E4	=	none	RIC1	26	296	1981	4309	7830
ES	1000000	none	RC1	26	985	3597	7371	10840
E6	11	none	RIC1	55	1062	3617	7464	10839
Amplification pa	Amplification performed with CT reagents, spi	T reagents, sp	iked with RIC1 primers (25 pmol RAN 16 and	mers (25 pmol 1	RAN 16 and 5 p	5 pmol T7/RAN 19	(6	
Each sample is	Each sample is an independent amplification.	amplification.						

FIG. 29